

**IN THE SPECIFICATION:**

**Please amend the paragraph beginning on page 24, line 16 as follows:**

~~Figure 26~~Figures 26A-26D when joined at match lines show~~show~~ a dendogram that illustrates the clustering relationship between deduced amino acid sequences of petunia (pCGP1907.aa), *Torenia* (pTMT5.aa) and *Fuchsia* (pCGP3267.aa) FMTs with other full length plant *O*-methyltransferases (OMT) of both Class I and Class II found in the GenBank database. The Genbank accession numbers of each SAM-OMT in the database are shown in brackets.